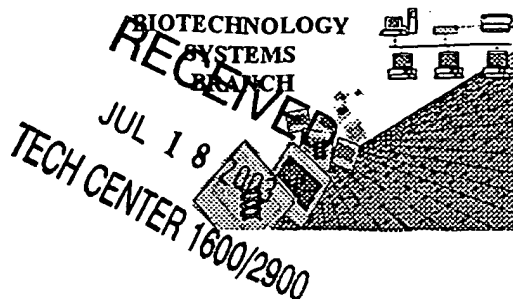


## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/939,226A  
Source: 1600  
Date Processed by STIC: 7/9/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003



1600

## RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

3 <110> APPLICANT: Euroscreen S.A.  
 4 SAMSON, Michel  
 5 PARMENTIER, Marc  
 6 VASSART, Gilbert  
 7 LIBERT, Frederick  
 9 <120> TITLE OF INVENTION: Methods for Identifying Compounds which Bind the Active CCR5  
 10 Chemokine Receptor  
 12 <130> FILE REFERENCE: 9409/2023C  
 14 <140> CURRENT APPLICATION NUMBER: 09/939,226A  
 15 <141> CURRENT FILING DATE: 2001-08-24  
 17 <150> PRIOR APPLICATION NUMBER: US 08/833,752  
 18 <151> PRIOR FILING DATE: 1997-04-09  
 20 <150> PRIOR APPLICATION NUMBER: US 09/626,939  
 21 <151> PRIOR FILING DATE: 2000-07-27  
 23 <150> PRIOR APPLICATION NUMBER: US 08/810,028  
 24 <151> PRIOR FILING DATE: 1997-03-04  
 26 <160> NUMBER OF SEQ ID NOS: 18  
 28 <170> SOFTWARE: PatentIn version 3.2  
 30 <210> SEQ ID NO: 1  
 31 <211> LENGTH: 792  
 32 <212> TYPE: DNA  
 33 <213> ORGANISM: Homo sapiens  
 35 <400> SEQUENCE: 1  
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 38 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa ttagacatc 120  
 40 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact 180  
 42 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggacaaga 240  
 44 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 300  
 46 aaaaaatcaa tgtgaagcaa atcgagccc gcctcctgcc tccgctctac tcaactggtgt 360  
 48 tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc 420  
 50 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc 480  
 52 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt 540  
 54 gtcaactctt gacagggtc tattttatag gcttcttctc tggaatcttc ttcacatcc 600  
 56 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcttta aaagccagga 660  
 58 cggtcacctt tgggggtgtg acaagtgtga tcaactgggt ggtggctgtg tttgcgtctc 720  
 60 tcccaggaat catctttacc agatctcaaa aagaaggctc tcattacacc tgcagctctc 780  
 62 attttccata ca  
 65 <210> SEQ ID NO: 2  
 66 <211> LENGTH: 1477  
 67 <212> TYPE: DNA  
 68 <213> ORGANISM: Homo sapiens  
 71 <220> FEATURE:  
 72 <221> NAME/KEY: misc\_feature

6-7  
 Does Not Comply  
 Corrected Diskette Needed

## RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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74 <223> OTHER INFORMATION: Any nucleotide
76 <220> FEATURE:
77 <221> NAME/KEY: misc_feature
78 <222> LOCATION: (1384)..(1385)
79 <223> OTHER INFORMATION: Any nucleotide
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84 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc      120
86 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact      180
88 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga      240
90 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc      300
92 aaaaaatcaa tgtgaagcaa atcgagccc gcctcctgcc tccgctctac tcaactggtgt      360
94 tcactcttgg ttttgtgggc aacatgctgg tcactcctcat cctgataaac tgcaaaaggc      420
96 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg ttttcccttc      480
98 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt      540
100 gtcaactctt gacagggctc tttttatag gcttcttctc tgggaatcttc ttcatcatcc      600
102 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcctta aaagccagga      660
104 cggtcacctt tggggtggtg acaagtgtga tcaactgggt ggtggctgtg tttgcgtctc      720
106 tcccaggaat catctttacc agatctcaaa aagaaggtct tcattacacc tgcagctctc      780
108 atttccata cagtcagtat caattctgga agaatttcca gacattaaag atagtcattct      840
110 tggggtcgtt cctgccgctg cttgtcatgg tcatctgcta ctcggaatc ctaaaaactc      900
112 tgcttcggtg tcgaaatgag aagaagaggc acagggtgtg gaggcttata ttcaccatca      960
114 tgattgttta ttttctcttc tgggctccct acaacattgt ccttctcctg aacaccttcc      1020
116 aggaattctt tggcctgaat aattgcagta gctctaacag gttggacca gctatgcagg      1080
118 tgacagagac tcttgggatg acgactgct gcatcaaccc catcatctat gcctttgtcg      1140
120 gggagaagtt cagaaactac ctcttagtct tcttccaaaa gcacattgcc aaacgcttct      1200
122 gcaaatgctg ttctattttc cagcaagagg ctcccagagc agcaagctca gtttacaccc      1260
124 gatccactgg ggagcaggaa atatctgtgg gcttgtgaca cggactcaag tgggctggtg      1320
W--> 126 acccagtcag agttgtgcac atggcttagt tttcatacac agcctgggct gggggtnggt      1380
128 tggnnagagt cttttttaaa aggaagttac tgttatagag ggtctaagat tcatccattt      1440
130 atttggcatc tgtttaaagt agattagatc cgaattc      1477
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134 <211> LENGTH: 1442
135 <212> TYPE: DNA
136 <213> ORGANISM: Homo sapiens
138 <400> SEQUENCE: 3
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141 tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc      120
143 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact      180
145 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga      240
147 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc      300
149 aaaaaatcaa tgtgaagcaa atcgagccc gcctcctgcc tccgctctac tcaactggtgt      360
151 tcactcttgg ttttgtgggc aacatgctgg tcactcctcat cctgataaac tgcaaaaggc      420
153 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg ttttcccttc      480
155 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt      540
157 gtcaactctt gacagggctc tttttatag gcttcttctc tgggaatcttc ttcatcatcc      600
159 tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcctta aaagccagga      660
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## RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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163 tcccaggaat catctttacc agatctcaaa aagaaggctc tcattacacc tgcagctctc 780
165 attttccata cattaaagat agtcatcttg gggctgggcc tgccgctgct tgtcatgggc 840
167 atctgctact cggaatcct aaaaactctg cttcgggtgc gaaatgagaa gaagaggcac 900
169 agggctgtga ggcttatctt caccatcatg attgtttatt ttctcttctg ggctccctac 960
171 aacattgtcc ttctcctgaa caccttccag gaattctttg gcctgaataa ttgcagtagc 1020
173 tctaacaggt tggaccaagc tatgcagggtg acagagactc ttgggatgac gcaactgtgc 1080
175 atcaacccca tcatctatgc ctttgtcggg gagaagtcca gaaactacct cttagtcttc 1140
177 ttccaaaagc acattgccaa acgcttctgc aaatgctgtt ctattttcca gcaagaggct 1200
179 cccgagcgag caagctcagt ttacaccgga tccactgggg agcaggaaat atctgtgggc 1260
181 ttgtgacacg gactcaagtg ggctgggtgac ccagtcagag ttgtgcacat ggcttagttt 1320
183 tcatacacag cctgggctgg ggggtggttg gaggtctttt ttaaaaggaa gttactgtta 1380
185 tagagggtct aagattcatc catttatttg gcatctgttt aaagtagatt agatccgaat 1440
187 tc 1442

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190 &lt;210&gt; SEQ ID NO: 4

191 &lt;211&gt; LENGTH: 184

192 &lt;212&gt; TYPE: PRT

193 &lt;213&gt; ORGANISM: Homo sapiens

195 &lt;400&gt; SEQUENCE: 4

```

197 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
198 1 5 10 15
201 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
202 20 25 30
205 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
206 35 40 45
209 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
210 50 55 60
213 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
214 65 70 75 80
217 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
218 85 90 95
221 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
222 100 105 110
225 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
226 115 120 125
229 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
230 130 135 140
233 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
234 145 150 155 160
237 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
238 165 170 175
241 Thr Cys Ser Ser His Phe Pro Tyr
242 180

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245 &lt;210&gt; SEQ ID NO: 5

246 &lt;211&gt; LENGTH: 352

247 &lt;212&gt; TYPE: PRT

248 &lt;213&gt; ORGANISM: Homo sapiens

250 &lt;400&gt; SEQUENCE: 5

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252 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
253 1 5 10 15

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## RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

```

256 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
257          20          25          30
260 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
261          35          40          45
264 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
265          50          55          60
268 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
269 65          70          75          80
272 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
273          85          90          95
276 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
277          100         105         110
280 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
281          115         120         125
284 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
285          130         135         140
288 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
289 145          150          155          160
292 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
293          165         170         175
296 Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn
297          180         185         190
300 Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu
301          195         200         205
304 Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys
305          210         215         220
308 Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile
309 225          230         235         240
312 Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu
313          245         250         255
316 Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser
317          260         265         270
320 Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr
321          275         280         285
324 His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe
325          290         295         300
328 Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe
329 305          310         315         320
332 Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser
333          325         330         335
336 Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu
337          340         345         350
340 <210> SEQ ID NO: 6
341 <211> LENGTH: 215
342 <212> TYPE: PRT
343 <213> ORGANISM: Homo sapiens
345 <400> SEQUENCE: 6
347 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
348 1          5          10          15

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## RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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351 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
352      20      25      30
355 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
356      35      40      45
359 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
360      50      55      60
363 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
364 65      70      75      80
367 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
368      85      90      95
371 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
372      100     105     110
375 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
376      115     120     125
379 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
380      130     135     140
383 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
384 145     150     155     160
387 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
388      165     170     175
391 Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala
392      180     185     190
395 Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Leu Gly Asn Pro Lys
396      195     200     205
399 Asn Ser Ala Ser Val Ser Lys
400      210     215
403 <210> SEQ ID NO: 7
404 <211> LENGTH: 360
405 <212> TYPE: PRT
406 <213> ORGANISM: Homo sapiens
409 <220> FEATURE:
410 <221> NAME/KEY: MISC_FEATURE
411 <222> LOCATION: (325)..(327)
412 <223> OTHER INFORMATION: Xaa = any amino acid
414 <400> SEQUENCE: 7
416 Met Leu Ser Thr Ser Arg Ser Arg Phe Ile Arg Asn Thr Asn Glu Ser
417 1      5      10      15
420 Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys
421      20      25      30
424 His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu
425      35      40      45
428 Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val
429      50      55      60
432 Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr
433 65      70      75      80
436 Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Ile Ile Thr Leu Pro
437      85      90      95
440 Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met
441      100     105     110

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<210> SEQ ID NO 14  
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<212> TYPE: DNA  
<213> ORGANISM: Artificial Sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: :  
<400> SEQUENCE: 14  
tcgaggatcc aagatggatt atcaagt

*see p.7 for error explanation*

27

<210> SEQ ID NO 15  
<211> LENGTH: 27  
<212> TYPE: DNA  
<213> ORGANISM: Artificial Sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: :  
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27

<210> SEQ ID NO 16  
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<212> TYPE: DNA  
<213> ORGANISM: Artificial Sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: :  
<400> SEQUENCE: 16  
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20

<210> SEQ ID NO 17  
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<212> TYPE: DNA  
<213> ORGANISM: Artificial Sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: :  
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27

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003  
TIME: 15:53:55

FYI

Input Set : A:\sequencelisting.txt  
Output Set: N:\CRF4\07092003\I939226A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1377,1384,1385  
Seq#:7; Xaa Pos. 325,326,327  
Seq#:8; Xaa Pos. 231,232,233,333,334,335  
Seq#:10; Xaa Pos. 145,146,147,321,322,323

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)  
(Sec.1.823 of new Rules)

Seq#:14,15,16,17

↑  
error explanation



VERIFICATION SUMMARY

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:55

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1320  
M:341 Repeated in SeqNo=2  
L:496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:320  
L:582 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:224  
M:341 Repeated in SeqNo=8  
L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:144  
M:341 Repeated in SeqNo=10  
L:885 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14  
L:887 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:14, <213>  
ORGANISM:Artificial Sequence  
L:887 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>  
ORGANISM:Artificial Sequence  
L:887 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:887  
L:894 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15  
L:896 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:15, <213>  
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L:912 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17  
L:914 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:17, <213>  
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L:914 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>  
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